

All Results
[S Melnik](#)
[E Rahm](#)
[R Durbin](#)
[A Woolfe](#)
[S Eddy](#)
[Comparison of Schema Matching Evaluations](#) - [all 13 versions »](#)

HH Do, S Melnik, E Rahm - Web, Web-Services, and Database Systems: Node 2002 Web-And 2003 - books.google.com
 ... sliveness thus preventing a qualitative **comparison** between systems. ... effort and the hardness of **match problems**. ... very helpful to better **compare** the effectiveness ...

[Cited by 144](#) - [Related Articles](#) - [Web Search](#)

[A computer vision based technique for 3-D sequence-independent structural comparison of proteins](#) - [all 3 versions »](#)

O Bachar, D Fischer, R Nussinov, H Wolfson - Protein Engineering Design and Selection, 2003 - Oxford Univ Press
 ... two typical structure **comparison** problems and **compare** them with ... provides an option to allow self **comparison**, which produces the best match besides die ...

[Cited by 75](#) - [Related Articles](#) - [Web Search](#)

[Large-Scale Comparison of Intron Positions among Animal, Plant, and Fungal Genes](#) - [all 5 versions »](#)

A Fedorov, AF Merican, W Gilbert - Proceedings of the National Academy of Sciences of the 2002 - National Acad Sciences
 ... alignment to **compare** these 30 genes to fungal-intron ... introns from the best fungal-animal **match** onto the ... such an animal-plant-fungal **comparison**, and Table ...

[Cited by 72](#) - [Related Articles](#) - [Web Search](#)

[\(PDF\) Hidden Markov Models in Computational Biology](#) - [all 18 versions »](#)

A Krogh, I Brown - J. Mol. Biol, 1994 - cab.zju.edu.cn
 ... of an HMM are estimated from a training **set** of unaligned sequences. ... To demonstrate the method, we examine three protein families. Each family consists of a **set** ...

[Cited by 644](#) - [Related Articles](#) - [View as HTML](#) - [Web Search](#) - [Library Search](#)

[YAP3: improved detection of similarities in computer program and other texts](#) - [all 7 versions »](#)

MJ Wise - Proceedings of the twenty-seventh SIGCSE technical symposium 1996 - portal.acm.org
 ... the problem exists of finding the initial **set** of unique ... to reduce the otherwise $O(n^2)$ cost of this **comparison**. ... of maximal-matches length L do **remove match**(p, t) ...

[Cited by 74](#) - [Related Articles](#) - [Web Search](#)

[Learning to match and cluster large high-dimensional data sets for data integration](#) - [all 10 versions »](#)

WW Cohen, J Richman - Proceedings of the eighth ACM SIGKDD International 2002 - portal.acm.org
 ... and • in Step 3d, let $T_{iaht} = 0$ (ie, only **remove** a from the **set** of PossibleCenters ...
 Two of the datasets require clustering, and two require **match-**ing ...

[Cited by 107](#) - [Related Articles](#) - [Web Search](#)

[S-Match: an Algorithm and an Implementation of Semantic Matching](#) - [all 16 versions »](#)

F Giunchiglia, P Shvaiko, M Yatskevich - The Semantic Web: Research and Applications: First European 2004 - books.google.com

... step 3) and, finally (step 4), we **compare** C_2 in ... Evaluation We have done some preliminary **comparison** between S ... is an estimate of the post match efforts needed ...

[Cited by 229](#) - [Related Articles](#) - [Web Search](#)

[Pastiche: making backup cheap and easy](#) - [all 11 versions »](#)

LP Cox, CD Murray, BD Noble - ACM SIGOPS Operating Systems Review. 2002 - portal.acm.org
 ... meta-data is **is not chunked** and is **mu- table**. ... with commonly-appearing chunkids, hoping for a false **match**. ... We call these the add **set**, delete **set**, and meta-data ...

[Cited by 150](#) - [Related Articles](#) - [Web Search](#)

[Fast algorithms for large-scale genome alignment and comparison](#) - [all 33 versions »](#)

AT Deicher, A Phillipcy, J Carlton, SL Salzberg - Nucleic Acids Research. 2002 - Oxford Univ Press
 ... $4k + 1$ bytes for a minimum **match** size of ... algorithmic extensions that permit alignment and **comparison** of protein ... It has been used to **compare** sequences as long ...

[Cited by 226](#) - [Related Articles](#) - [Web Search](#)

[\[PDF\] Automatic grammar induction and parsing free text: A transformation-based approach](#) - [all 20 versions](#)

E Brill - Proceedings of the 31st Annual Meeting of the Association ... 1993 - [acl.ldc.upenn.edu](#)

... presented above allow us to **compare** the transformation ... carried out are integer addition and integer **comparison**. ... hope that this grammar will **match** the grammar ...

[Cited by 182](#) - [Related Articles](#) - [View as HTML](#) - [Web Search](#)



Result Page: [1](#) [2](#) [3](#) [4](#) [5](#) [6](#) [7](#) [8](#) [9](#) [10](#) [Next](#)

[Google Home](#) - [About Google](#) - [About Google Scholar](#)

©2008 Google